

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/900,519A

CRF Processing Date: 3/25/2002  
 Edited by: [Signature]  
 Verified by: [Signature] (STIC staff)

**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



OIPE

## RAW SEQUENCE LISTING

DATE: 03/25/2002

PATENT APPLICATION: US/09/900,519A

TIME: 12:41:40

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\03222002\I900519A.raw

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4 <110> APPLICANT: Allen, Keith D.
6 <120> TITLE OF INVENTION: TRANSGENIC MICE CONTAINING
7   ADRENOMEDULLIN RECEPTOR GENE DISRUPTIONS
10 <130> FILE REFERENCE: R-615
12 <140> CURRENT APPLICATION NUMBER: US 09/900,519A
13 <141> CURRENT FILING DATE: 2001-07-06
15 <150> PRIOR APPLICATION NUMBER: US 60/216,254
16 <151> PRIOR FILING DATE: 2000-07-06
18 <150> PRIOR APPLICATION NUMBER: US 60/221,497
19 <151> PRIOR FILING DATE: 2000-07-27
21 <150> PRIOR APPLICATION NUMBER: US 60/280,264
22 <151> PRIOR FILING DATE: 2001-03-29
24 <160> NUMBER OF SEQ ID NOS: 3
26 <170> SOFTWARE: FastSEQ for Windows Version 4.0
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 526
30 <212> TYPE: DNA
31 <213> ORGANISM: Mus spretus
33 <400> SEQUENCE: 1
34 tccatgctgc actgtgtggc caaccccatc ctctacaact ttctcagccc gagcttccgg 60
35 ggccgactgc tgagccttgt ggttcgttac cttcccaagg agcaggccag ggcagcaggt 120
36 ggtcgtgcct cctcttcttc ttccaccag cactccatca tcattaccaa agagggcagc 180
37 ctgcccgcgtg cagcggatct ccacaccac cccatccgaa acgttcaggc gtcctctccg 240
38 cctccaaaca cctcacctac actctgcaat tccgtagcca gctaaggtag actctagctt 300
39 cctccaccaa taagaaagtt cagaggggga tgcgagaggt ctgtgggagg ggggtgggaag 360
40 gactggcttg ttcagggccca atttaagtat atcaaaacgt tgctgtgggg agagggaaac 420
41 ggttcgggaa ggacagagaa tggatctttc cttgatagta cactatttgt ttgggtactg 480
42 atgtctaagg gagccacacc ggtggggcgt ggggggtggg gaagcg 526
44 <210> SEQ ID NO: 2
45 <211> LENGTH: 200
46 <212> TYPE: DNA
47 <213> ORGANISM: Artificial Sequence
49 <220> FEATURE:
50 <223> OTHER INFORMATION: Targeting Vector
52 <400> SEQUENCE: 2
53 ggcgccactg cctgttgatg tgggcttaca tagttgtctt tgccatctgc tggctgccct 60
54 accaagtgc tatgctgctg ctactctgc acgggaccca catcttcctc cactgtcacc 120
55 tggtaacct tctctacttc ttctacgaaa tcatcgactg cttttccatg ctgcactgtg 180
56 tggccaacct catcctctac 200
58 <210> SEQ ID NO: 3
59 <211> LENGTH: 200
60 <212> TYPE: DNA
61 <213> ORGANISM: Artificial Sequence

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VERIFICATION SUMMARY

DATE: 03/25/2002

PATENT APPLICATION: US/09/900,519A

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